

FIG. 2

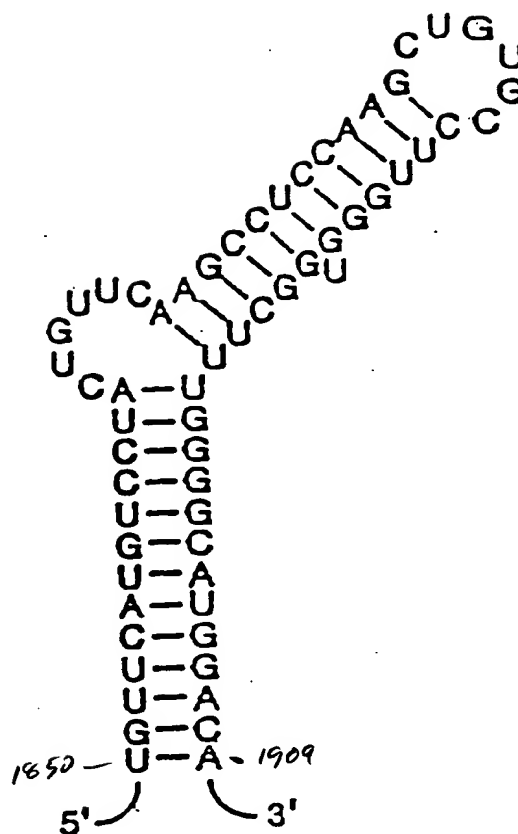


FIG. 3

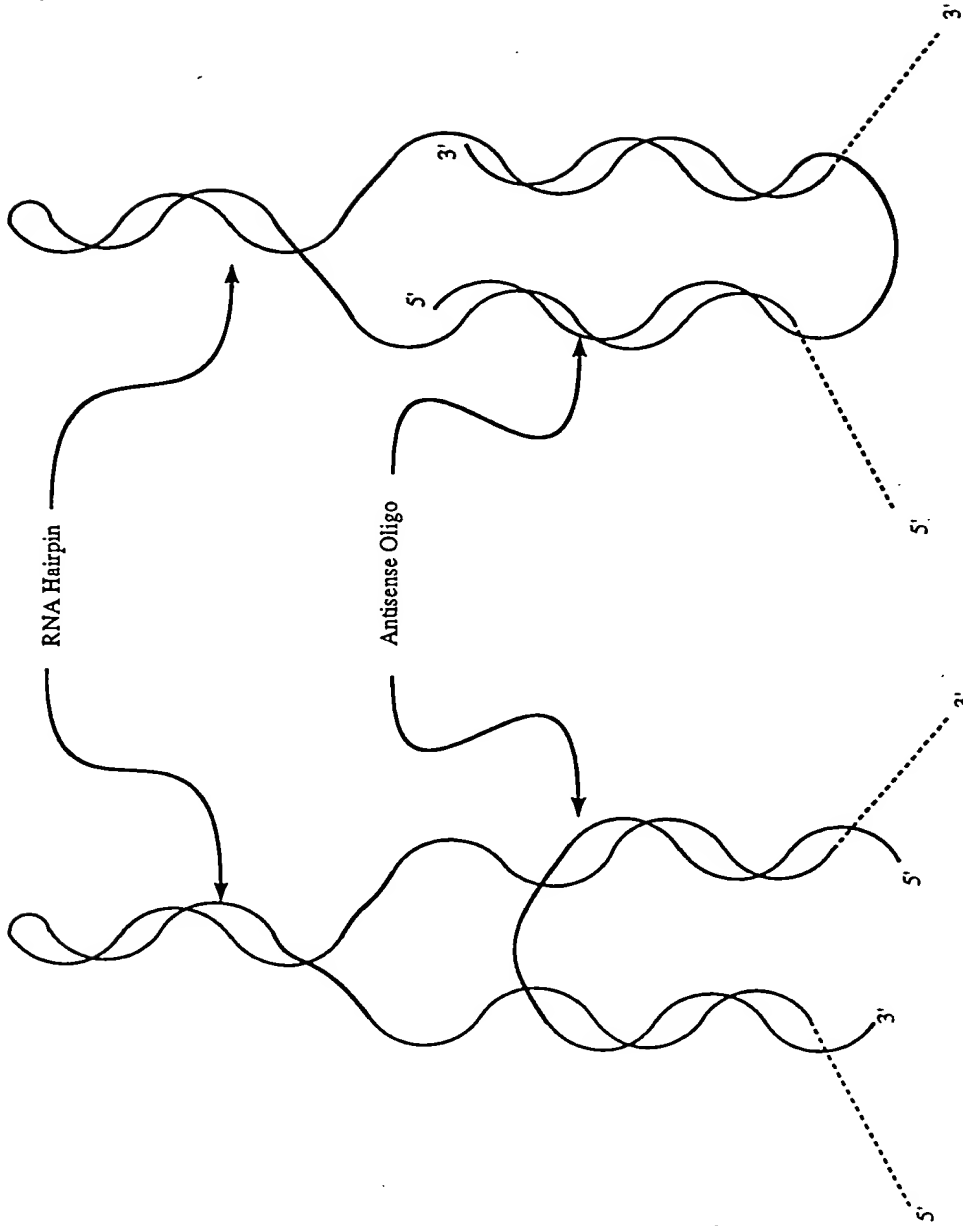


FIG. 4B

FIG. 4A

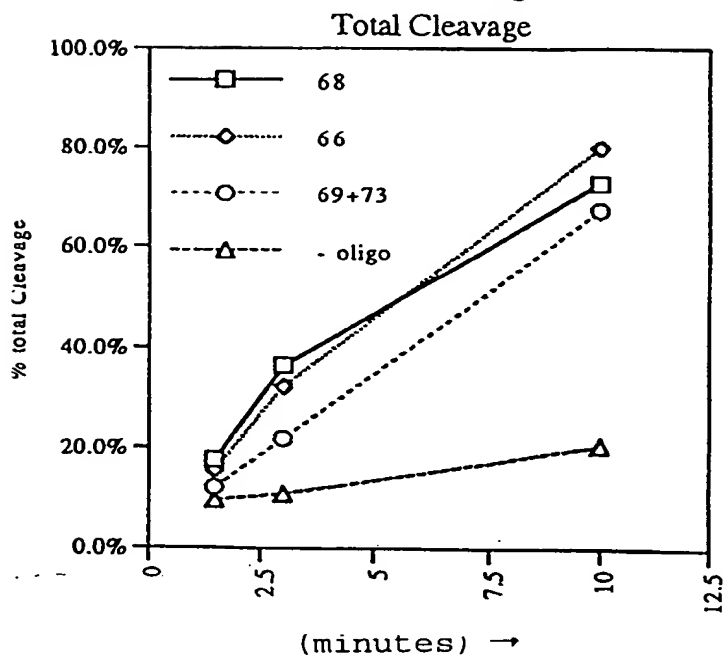


FIG. 5A

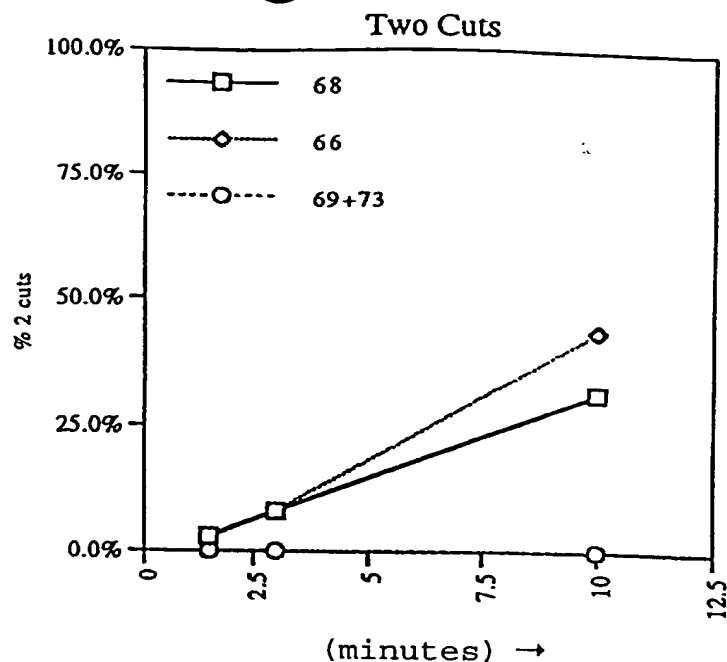


FIG. 5B

RNase H Cleavage with 12+12 Noncontiguous Phosphorothioates

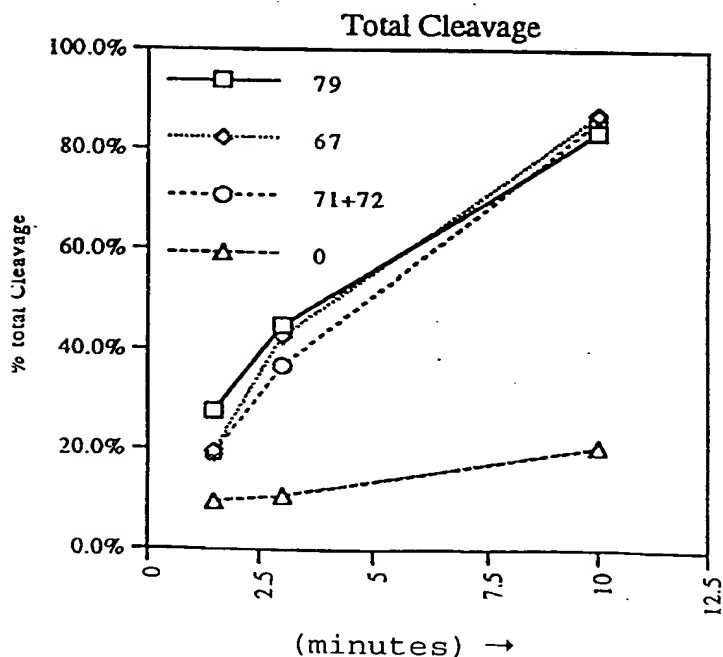


FIG. 6A

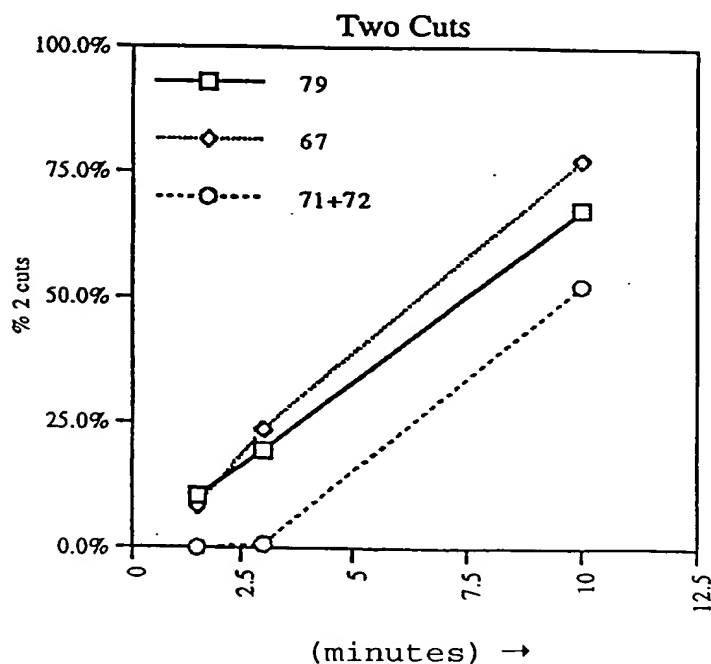


FIG. 6B

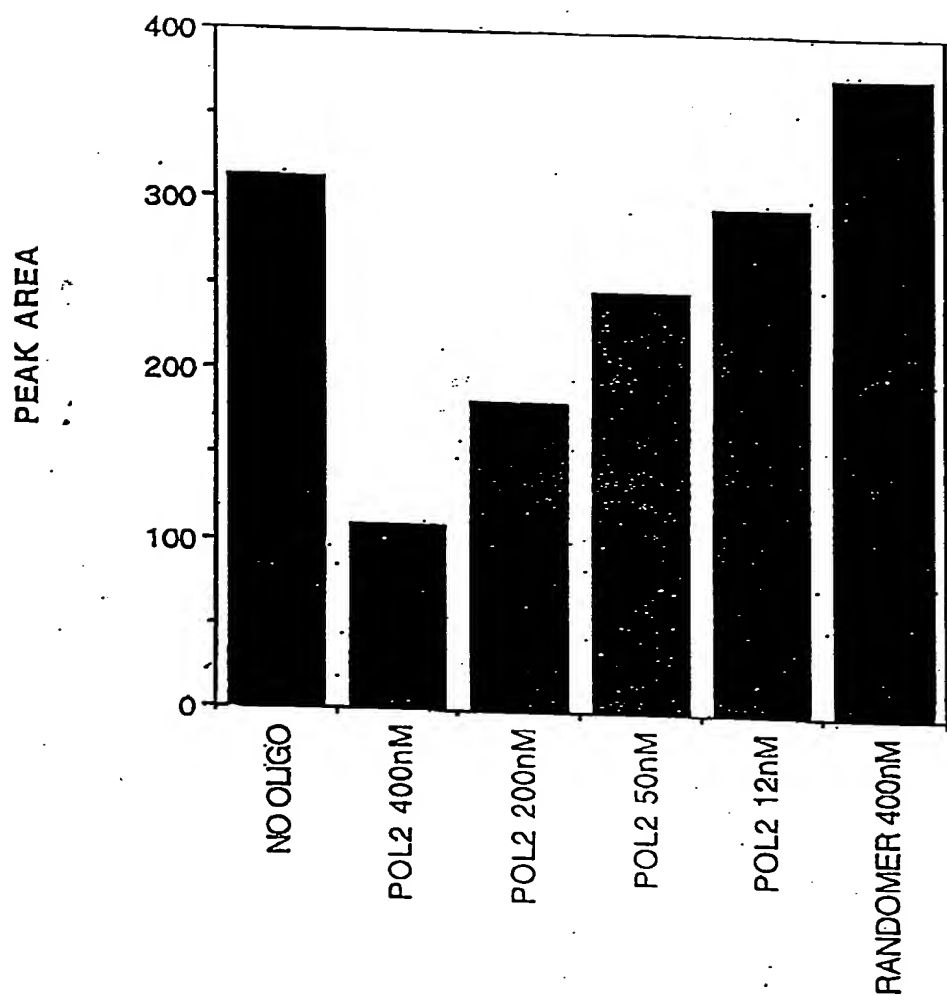
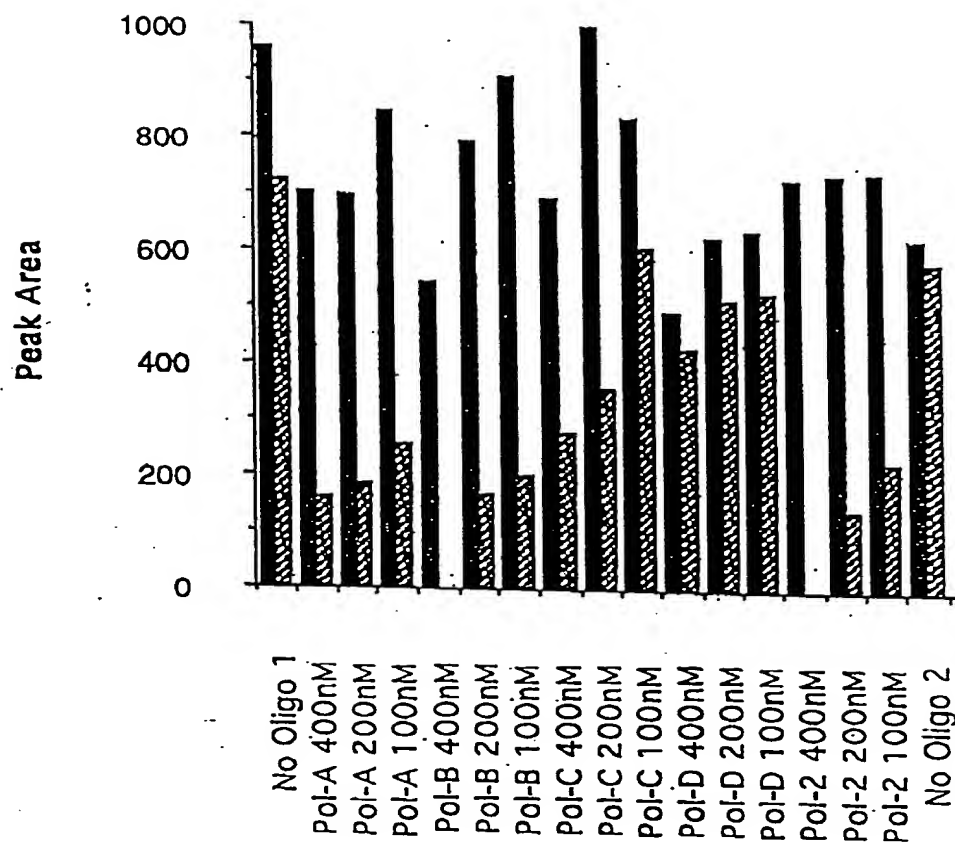


FIG. 7



Dark bars represent translation of HSVProt control RNA
Hatched bars represent translation of HBVPol test RNA

FIG. 8

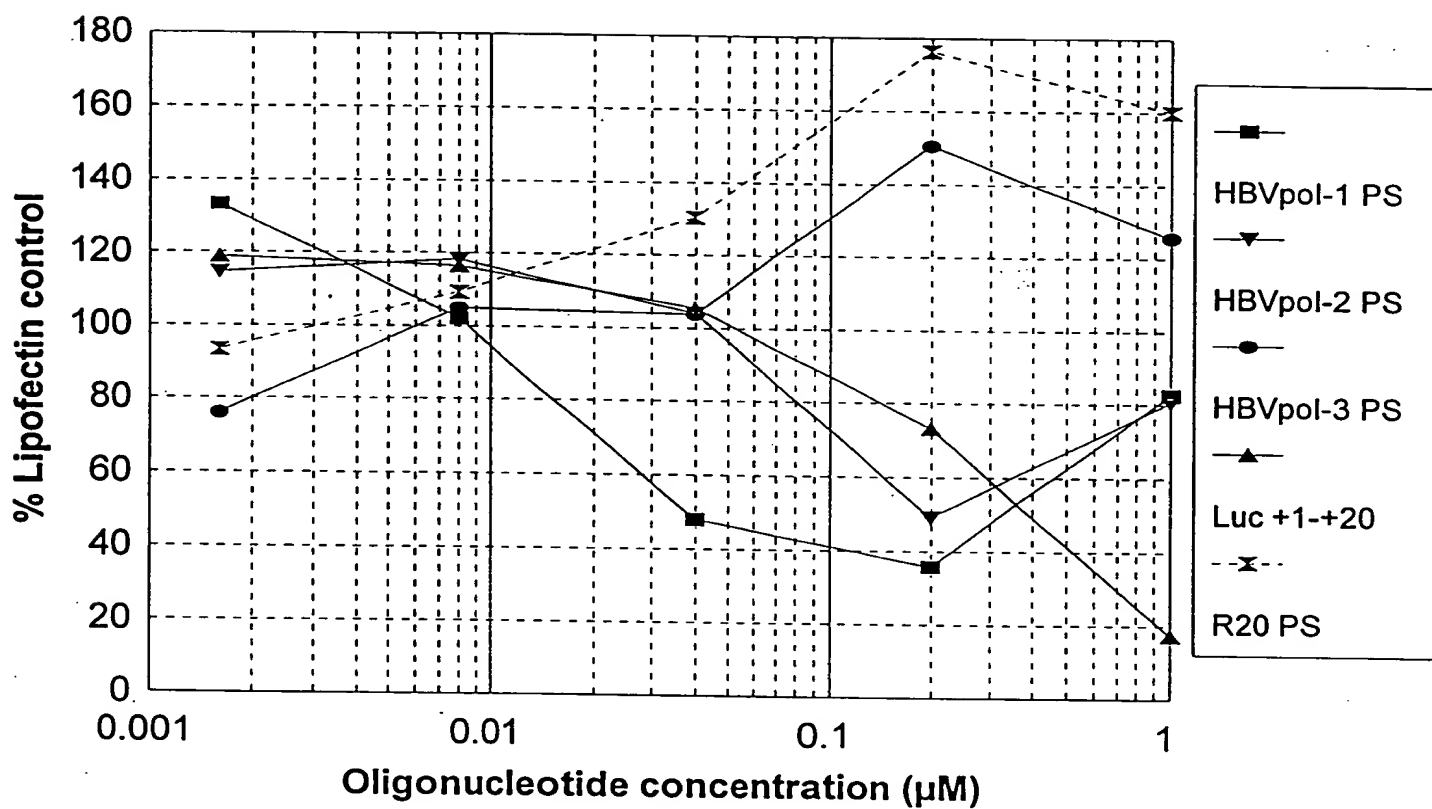
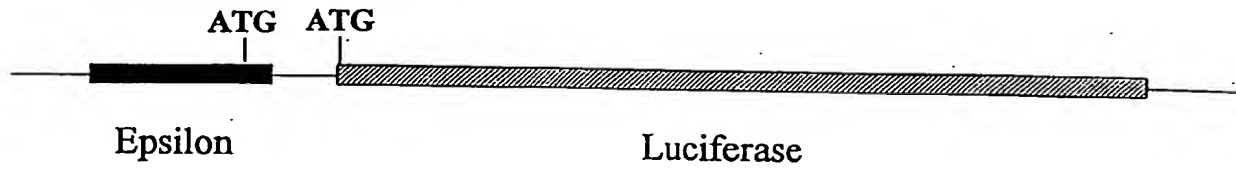


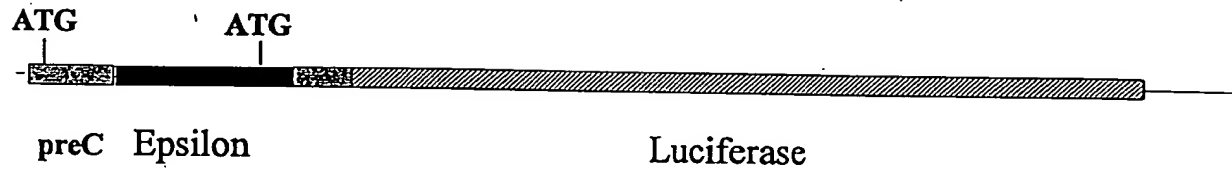
FIG. 9

Alternate HBV epsilon targets for primary assays

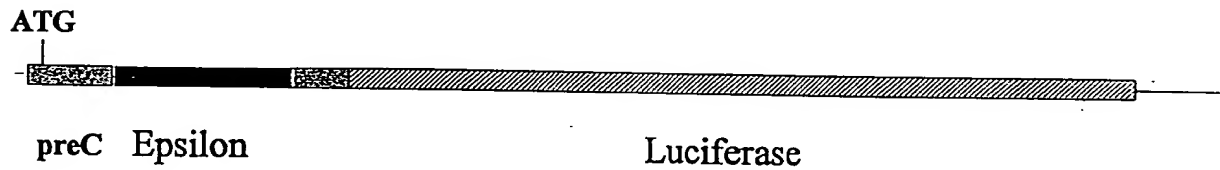
pGLE



pGLE2



pGLE3



pGL_{pol}

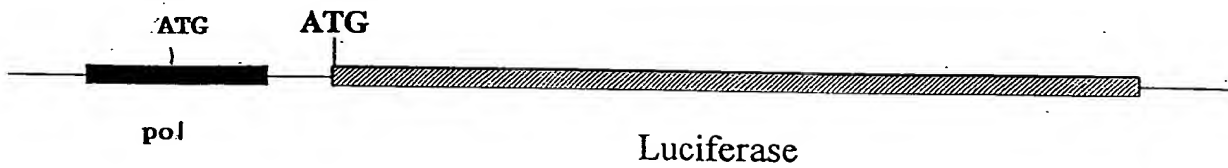


FIG. 10

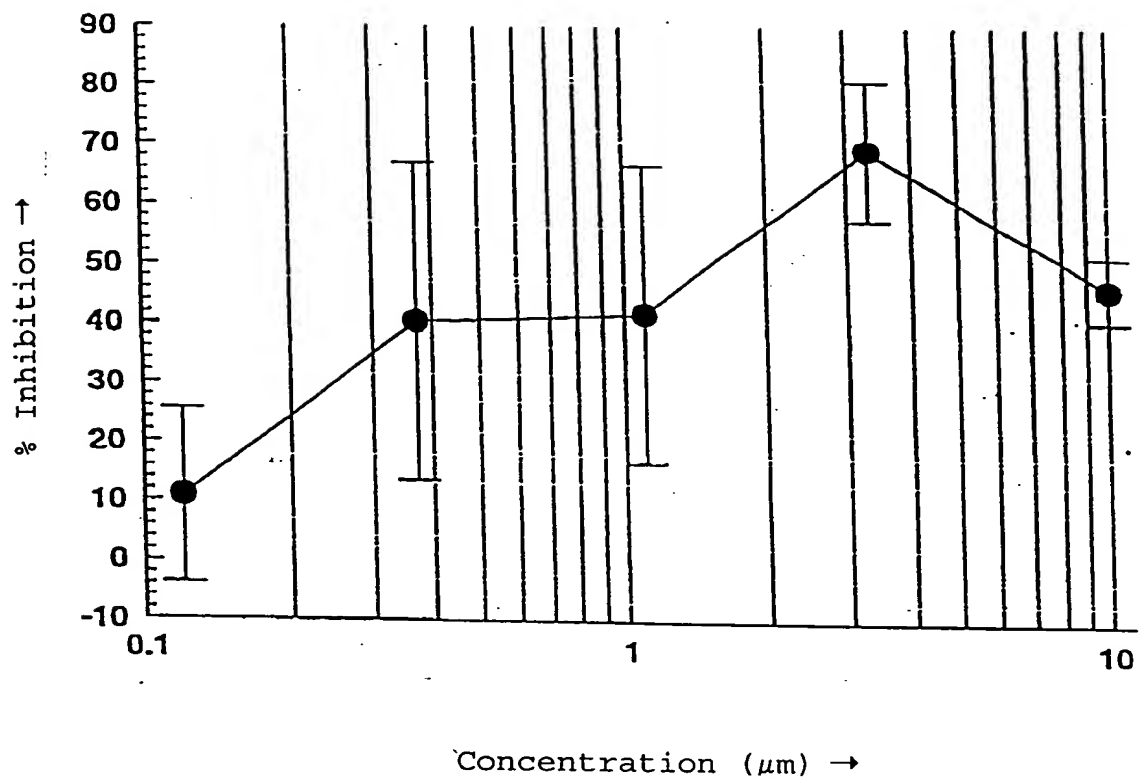


FIG. 11

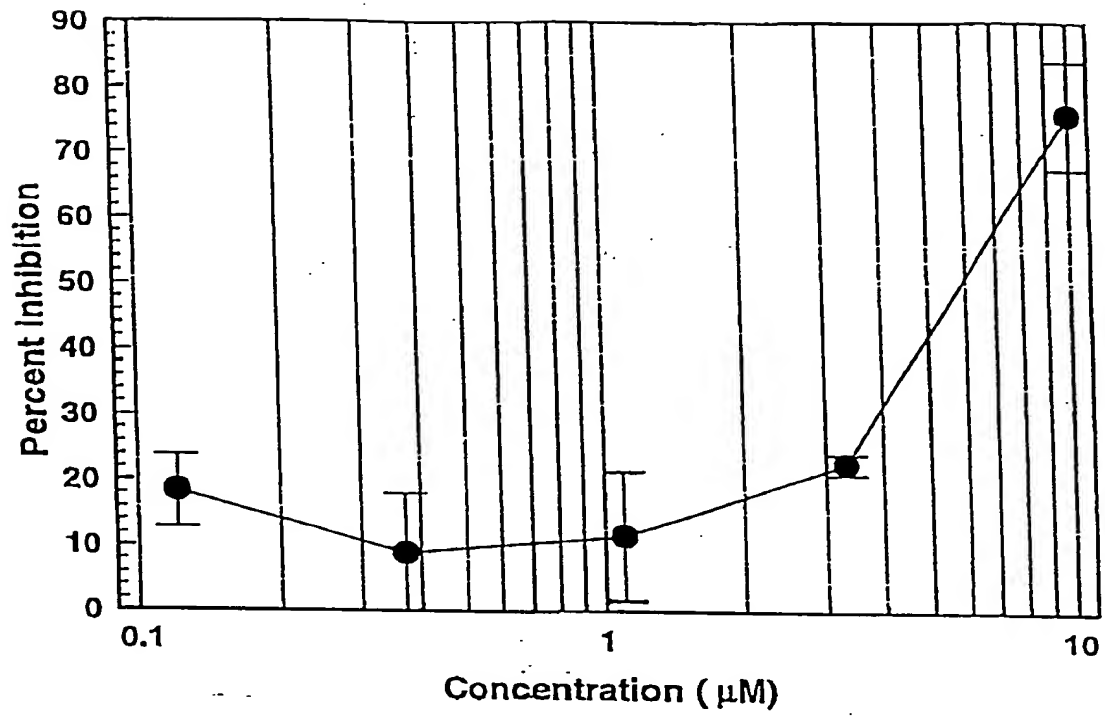
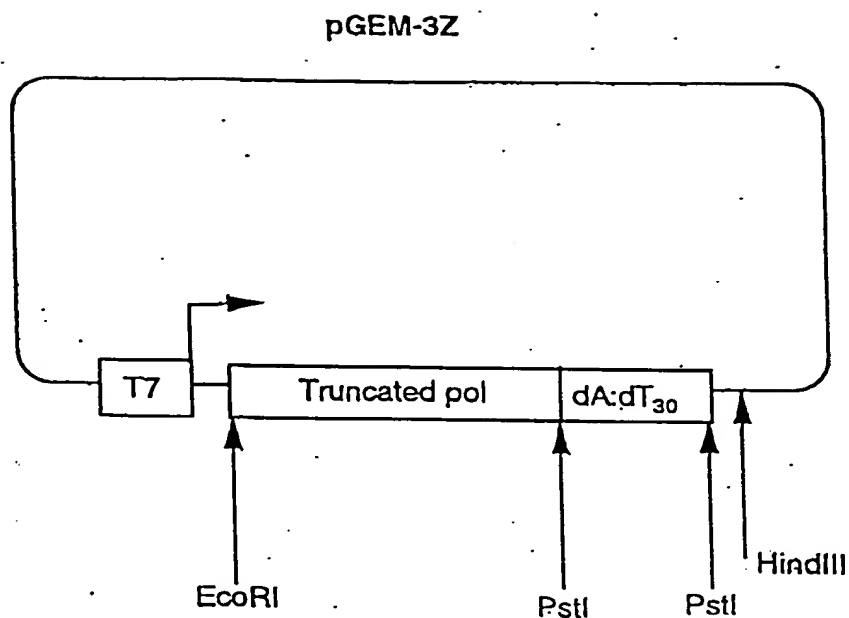


FIG. 12



Truncated pol represents -17 to +634 of the pol open reading frame. The normal leucine codon at +635 to +637 has been replaced by a stop codon.

FIG. 13